



## RAW SEQUENCE LISTING ERROR REPORT

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
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U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
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Revised 01/29/2002

10/03/33/A

SEQUENCE LISTING

<110> JAPAN SCIENCE AND TECHNOLOGY CORPORATION

<120> Screening of genes to give tolerance against environmental stress and the applications

<130> 12-130

"1" applications

<140>

<141>

<150> JP P1999-235910

<151> 1999-07-19

<150> JP P2000-85377

<151> 2000-03-24

<160> 66

<170> PatentIn Ver. 2.1

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<220>

<221> CDS

<222> (42)..(464)

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Arg Ser Lys Ser Ala Thr Val Pro Thr Gln Thr Val Leu Pro Phe Lys  
25 30 35

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Arg Ser Gln Ala Thr Pro Ser Asp Asp Ala Ser Arg Pro Thr Lys Val  
70 75 80 85

caa gag ctg tgt gtg tat gag aac gag aga gat cgt gga agc cct 344

Does Not Comply  
Corrected Diskette Needed

( see p.2 )

( only the  
beginning and the  
end of first file  
shown )

10/031,331 A 2

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Gln Ser Val Glu Glu Ser Lys Gly Tyr Gly Gly His Gly Gly His

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end of first file

Do NOT divide the Sequence Listing  
into separate files.

Per 1.824 of Sequence Rules, "The computer readable  
form shall contain a single 'Sequence Listing'..."

Submit one Sequence Listing file containing  
all the sequences.

Use of n and/or Xaa has been detected in the Sequence Listing.  
Review the Sequence Listing to insure a corresponding  
explanation is presented in the <220> to <223> fields of  
each sequence using n or Xaa.

FJF